

Epidemiology and pathogenesis of Kaposi's sarcoma-associated herpesvirus

Chris Boshoff^{1*} and Robin A. Weiss²

¹Department of Oncology and Molecular Pathology, The Wolfson Institute for Biomedical Research, Cruciform Building,
University College London, Gower Street, London WC1 6BT, UK

²Department of Immunology and Molecular Pathology, Wohl Virion Centre, Windeyer Institute of Medical Sciences,
46 Cleveland Street, London W1T 4JF, UK

Kaposi's sarcoma (KS) occurs in Europe and the Mediterranean countries (classic KS) and Africa (endemic KS), immunosuppressed patients (iatrogenic or post-transplant KS) and those with acquired immune deficiency syndrome (AIDS), especially among those who acquired human immunodeficiency virus sexually (AIDS–KS). KS-associated herpesvirus (KSHV or HHV-8) is unusual among herpesviruses in having a restricted geographical distribution. Like KS, which it induces in immunosuppressed or elderly people, the virus is prevalent in Africa, in Mediterranean countries, among Jews and Arabs and certain Amerindians. Distinct KSHV genotypes occur in different parts of the world, but have not been identified as having a differential pathogenesis. KSHV is aetiologically linked to three distinct neoplasms: (i) KS, (ii) primary effusion lymphoma, and (iii) plasmablastic multicentric Castleman's disease. The histogenesis, clonality and pathology of the tumours are described, together with the epidemiology and possible modes of transmission of the virus.

Keywords: Kaposi's sarcoma-associated herpesvirus; human herpesvirus 8; Kaposi's sarcoma; body cavity-based lymphoma; primary effusion lymphoma; Castleman's disease

1. EPIDEMIOLOGY OF KAPOSI'S SARCOMA

Nearly 40 million people worldwide are now infected by the human immunodeficiency virus (HIV-1), not counting the 20 million people who have already died from HIV-related diseases. In Africa, where most of these people live, the vast majority of HIV-infected individuals still succumb to infectious diseases, most notably tuberculosis. However, opportunistic neoplasms are also taking their toll and Kaposi's sarcoma, for example, is now the most common tumour in men in Uganda (Ziegler et al. 1997).

In 1872, the Hungarian dermatologist Moritz Kaposi published the case histories of five patients with 'idiopathic multiple pigmented sarcomas' of the skin (Kaposi 1872). This form of the disease was eponymously designated Kaposi's sarcoma (KS) in 1891. Kaposi's original patients had a more aggressive disease than HIV-negative cases in Europe today, with tumours described in the pharynx, stomach, small intestine, liver and colon in addition to the skin (Kaposi 1872). It is not known whether these patients were significantly immunosuppressed due to a concurrent illness. Four epidemiological forms of KS are currently distinguished: classic KS in Europe and the Mediterranean; endemic, HIV-negative KS in Africa; post-transplant or iatrogenic KS; and HIV-associated KS.

Classic KS occurs predominantly in elderly patients of southern European, Arabic or Jewish ancestry (Franceschi & Geddes 1995), where it usually remains as an indolent

*Author for correspondence (c.boshoff@ucl.ac.uk).

disease affecting the extremities. It is more common in men than women (sex ratios are estimated to be from 3:1 to 10:1) (Franceschi & Geddes 1995). Within countries where classic KS occurs there is also a specific geographical distribution; e.g. in Greece KS occurs more commonly in those from the Peloponnese peninsula. In Italy, higher incidence rates are reported in Sicily and southern Italy than in northern Italy (Bertaccini 1959; De Amicis 1897). Between 1977 and 1991 the annual incidence of classic KS in Sardinia per 100 000 inhabitants was 2.43 for men and 0.77 for women (ratio 3.2:1) (Cottoni et al. 1996). In eastern European countries KS predominates in those of Ashkenazi Jewish heritage (Rothman 1962).

In some equatorial countries of Africa, KS has existed for many decades, pre-dating the emergence of HIV. This form of the disease is known as endemic KS and is usually more aggressive than the classic form (D'Oliveira & Torres 1972). Lymph nodes are often affected and the disease is seen in both children and adults (Olweny et al. 1976; Ziegler & Katongole-Mbidde 1996). Whereas the median age of classic KS is 65 years, the median age of endemic KS is 40 years (Oettle 1962). As reports of African endemic KS accumulated during the 1960s and 1970s it became clear that there is also a specific geographical distribution in Africa: KS is more common in equatorial Africa; e.g. Oettle (1962) reported that KS represented 10% of all malignancies in Uganda with a male to female ratio in adults of 15:1.

During the past three decades, the incidence of KS among renal transplant recipients and other patients

Table 1. KS incidence from population-based registries

country	period	male	female	reference
classic KS ^a				
USA	1970-1979	0.29	0.10	Biggar <i>et al.</i> (1984)
UK	1975-1988	0.01	0.003	Grulich et al. (1992)
Israel	1970-1989	1.93	0.73	Iscovich et al. (1998)
Sardinia	1997-1991	2.43	0.77	Cottoni et al. (1996)
Africa in the AIDS erab				
Zimbabwe	1990-1992	23.6	10.1	Bassett <i>et al.</i> (1995)
Uganda	1991-1993	48.2^{c}	23.8	Wabinga <i>et al.</i> (1993)
Rwanda	1991-1994	9.5	2.5	Newton <i>et al.</i> (1996)
Guinea	1992-1995	0.2	0.2	Koulibaly et al. (1997)

^a KS incidence per 100 000 population, age-standardized.

receiving immunosuppressive therapy has increased (post-transplant or iatrogenic KS) (Harwood et al. 1979). The incidence of KS among renal transplant recipients is estimated to be approximately 150 times that seen in a healthy Western population (Penn 1983). In some countries where endemic or classic KS occurs (e.g. Saudi Arabia), KS is the most common post-transplant tumour (Qunibi et al. 1988). This again indicates that specific genetic or environmental factors are involved in its pathogenesis. In Europe, a retrospective study (1968–1990) among 7923 organ-transplant recipients recorded an overall prevalence of KS of 0.52% and KS was significantly more frequent after a liver (1.24%) than a kidney (0.45%) or heart (0.41%) transplant (Farge 1993). Patients of Mediterranean, Jewish or Arabian ancestry are also overrepresented among iatrogenic immunosuppressed patients who develop KS in North America (Franceschi & Serraino 1995). This indicates that those who were born or whose family originated from countries where classic KS occurs continue to be at higher risk of developing KS, even after migration to so called 'lowrisk' countries.

In 1981 the first cases of KS in young men from New York City and San Francisco were described (US Public Health Service 1981) heralding the beginning of the acquired immune deficiency syndrome (AIDS). KS is now the most common neoplasm occurring in patients with AIDS (Beral et al. 1990; Rabkin et al. 1995). Like the other forms of KS, AIDS-KS is not distributed randomly among HIV risk groups and surveillance data have consistently supported the existence of a sexually transmissible KS cofactor: in the West, AIDS-KS occurs predominantly in gay men with HIV, less commonly in those who acquired HIV through heterosexual contact and only rarely in patients who acquired HIV parenterally (patients with haemophilia or intravenous drug users) (Beral et al. 1990). In Africa, HIV infection has had an immense effect on the incidence of KS and the vast majority of patients with KS live on that continent. Recent reports from sub-Saharan Africa have confirmed the wide differences in incidence rates between countries and also between sexes (table 1).

In Uganda, among HIV-positive subjects, KS cases are characterized by better education and more affluence,

compared with controls (Ziegler et al. 1997). Urban, rather than rural address, exposure to water, travel away from home and sexually transmitted diseases are also more frequent among those with KS (Ziegler et al. 1997). These data may indicate that poverty protects against KS development. It has been proposed that early childhood infection, with an agent such as KS-associated herpesvirus (KSHV), confers long-lasting immunity, whereas acquisition of such an agent as sexually active adults would be less protective, especially with immunosuppression due to HIV (Ziegler et al. 1997).

2. KAPOSI'S SARCOMA HISTOGENESIS

Histologically, KS is a complex lesion: in early lesions there are a collection of irregular endothelial lined spaces that surround normal dermal blood vessels and these are accompanied by a variable inflammatory infiltrate (patch stage). This stage is followed by the expansion of a spindle-celled vascular process throughout the dermis. These spindle cells form slit-like, vascular channels containing erythrocytes (plaque stage). The later nodular stage KS lesions are composed of sheets of spindle cells, some of which are undergoing mitosis, and slit-like vascular spaces with areas of haemosiderin pigmentation. The spindle cells form the bulk of established KS lesions and are therefore thought to be the neoplastic component. Most of the spindle cells in KS lesions express endothelial markers, including CD31 and CD34. However, it was also shown that KS spindle cells express markers for smooth muscle cells, macrophages and dendritic cells (Nickoloff & Griffiths 1989; Sturzl et al. 1992) suggesting that spindle cells are either derived from pluripotent mesenchymal precursors or represent a heterogeneous population of cells. Recently, it was shown that all KS spindle cells express vascular endothelial growth factor receptor (VEGFR)-3 (Dupin et al. 1999; Jussila et al. 1998). VEGFR-3 is usually expressed only by lymphatic endothelium and by neoangiogenic vessels, but not by mature vascular endothelial cells, indicating that KS spindle cells probably belong to the endothelial lineage that can differentiate into lymphatic cells.

Endothelial stem cells have not yet been identified (Risau 1997), but putative endothelial progenitors

^b Percentage of all recorded cancers that are KS.

^c Although this study suggests that nearly 50% of all male cancers in Uganda are due to KS, there could be a bias towards HIV-infected patients attending central hospitals.

(angioblasts) involved in angiogenesis have been isolated from peripheral blood (Asahara et al. 1997). These cells express CD34 and the VEGF receptor Flk-1 and might be related to KS spindle cells. Circulating KS-like spindle cells have been isolated and cultured from patients with AIDS-KS and HIV-infected individuals thought to be at risk of developing KS. These circulating cells have an adherent phenotype and express markers for both macrophages and endothelial cells, as well as in vitro functional activities similar to KS spindle cells (Browning et al. 1994; Sirianni *et al.* 1997).

Tumours produce cytokines and their cells respond positively or negatively to cytokines in culture. KS is no exception and KS spindle cells or infiltrating CD8+ lymphocytes and macrophages express high levels of interleukin-6 (IL-6), basic fibroblast growth factor (bFGF), tumour necrosis factor-α, oncostatin M and γinterferon (IFN-γ) (Ensoli et al. 1989, 1994; Fiorelli et al. 1998; Miles et al. 1990; Nair et al. 1992; Salahuddin et al. 1988; Samaniego et al. 1995). IL-6 is produced by KS spindle cells and exogenous IL-6 enhances the proliferation of KS cells in culture (Miles et al. 1990). IFN-γ also induces endothelial cells to acquire similar phenotypic features to KS spindle cells (Fiorelli et al. 1998). Because of the nature of KS lesions it has been suggested that these tumours are 'cytokine driven'.

The more aggressive nature of HIV-associated KS has led to speculation that HIV-1-encoded proteins may directly enhance KS growth (Ensoli et al. 1994). The HIV-1 Tat protein transactivates HIV viral genes and also some host cell genes (Vaishnaw & Wong-Staal 1991). Tat can be released by infected cells and can act extracellularly (Ensoli et al. 1993; Frankel & Pabo 1988). Tat induces a functional programme in endothelial cells related to angiogenesis and inflammation including the migration, proliferation and expression of plasminogen activator inhibitor-1 and E selectin (Albini et al. 1995). Tat induces growth of KS spindle cells in vitro and is angiogenic in vivo and in transgenic mice (Ensoli et al. 1993, 1994; Vogel et al. 1988). The Tat basic domain contains an arginine- and lysine-rich sequence which is similar to that of other potent angiogenic growth factors including vascular endothelial growth factor-A (VEGF-A) and bFGF (Albini et al. 1996). Tat specifically binds to and activates the Flk-1/kinase domain receptor (Flk-1/KDR), a VEGF-A tyrosine kinase receptor (Albini et al. 1997). Tat-induced angiogenesis can be inhibited by agents blocking this receptor (Albini et al. 1997). The RGDcontaining region of Tat has also been postulated to have a role in the pathogenesis of AIDS-KS; however, baboons infected with HIV-2, whose Tat lacks an RGD sequence, can develop 'KS-like lesions', albeit with myofibroblast, rather than endothelial, phenotype (Barnett et al. 1994; Ensoli et al. 1994). AIDS-associated KS has a specific tissue distribution (often affecting the nose, oral and genital mucosae) and it is possible that the angiogenic or tumour-growth-promoting properties of Tat (Prakash et al. 2000) contribute to these phenomena.

The fact that the incidence of KS is higher in African countries where HIV-1 is prevalent, compared with countries where HIV-2 is more common, despite similar prevalences of antibodies to KSHV (Ariyoshi et al. 1998), has argued in favour of functional differences between HIV-1 and HIV-2 Tat proteins (Gallo 1998). Although the increased incidence in HIV-1-infected individuals of various neoplasia including KS, lymphoma and squamous carcinoma is well documented (Parkin et al. 1999), we are not aware of definitive cancer statistics from HIV-2-prevalent regions. HIV-1 Tat could indirectly affect tumour cell proliferation via the induction of cellular growth and angiogenic factors and/or directly activate transcriptional regulators like nuclear factor kappa B.

3. KAPOSI'S SARCOMA CLONALITY

The nature of KS also remains controversial as to whether it is a neoplastic lesion or a reactive process. Like Hodgkin's disease, the exact tumour cell type can be argued and the 'tumour cell' compartment in early lesions makes up the minority of the tumour bulk, where the majority of cells are inflammatory cells. Furthermore, the clinical presentation of multiple skin lesions in a defined distribution and the spontaneous remission of lesions favour a reactive hyperplasia rather than a true malig-

A useful marker for clonality is the inactivation pattern of X chromosomes in females with cancer (Vogelstein et al. 1985). The human X-linked androgen receptor gene can be used to assess clonal patterns of X chromosome inactivation, owing to common polymorphisms which can be identified by methylation-dependent DNA restriction enzymes. Using this technique, Rabkin et al. (1995) showed that individual KS lesions are probably clonal. However, a French group led by Oksenhendler demonstrated in skin lesions, including four with nodular KS where more than 70% of the cells were spindle cells, a polyclonal pattern of inactivation (Delabesse et al. 1997). Two other studies have indicated that multiple lesions in the same patient can be oligo- or monoclonal (Gill et al. 1998; Rabkin et al. 1997), suggesting that KS is a disseminated monoclonal cancer and that the changes that permit the clonal outgrowth of spindle cells occur before metastasis. The circulating cells giving rise to multiple clonal lesions are potentially related to the spindleshaped cells that can be cultured from peripheral blood, which are increased in HIV-infected patients who have KS or are at high risk of developing KS (Browning et al. 1994).

Judde et al. (2000) expanded a study first explored by the discoverers of KSHV (Russo et al. 1996) investigating KSHV clonality within tumours by analysing the size heterogeneity of KSHV-fused terminal repeats (TRs). This assay has been previously used to demonstrate that certain Epstein-Barr virus (EBV)-associated tumours are monoclonal (Raab-Traub & Flynn 1986). However, the TR region of KSHV is much larger, and rearrangements might therefore occur, skewing data towards oligo- or polyclonality. By using this technique Judde et al. (2000) showed that KSHV is monoclonal within nodular KS lesions, indicating that the virus was present prior to the expansion of a tumour clone of KS spindle cells.

Early KS (patch stage) is probably a non-clonal proliferation of lymphatic endothelial cells or endothelial precursors (e.g. angioblasts) (Risau 1997) with a prominent inflammatory and angiogenic response, whereas advanced disease can develop into a true clonal malignancy with

520

metastases of clonally derived spindle cells (Delabesse *et al.* 1997; Gill *et al.* 1998; Rabkin *et al.* 1995, 1997).

4. MOLECULAR DETECTION OF KAPOSI'S SARCOMA-ASSOCIATED HERPESVIRUS

A viral aetiology for KS was suspected long before the onset of the AIDS epidemic (Oettle 1962). In 1972, herpesvirus-like particles were found by electron microscopy in KS tumour cells and were shown to be cytomegalovirus (CMV) (Giraldo et al. 1972, 1975). DNA sequences of CMV, human herpesvirus 6 (HHV-6), human papilloma viruses (HPVs) and BK virus (human polyoma virus) and other viral or bacterial pathogens have all been detected in KS lesions and put forward as suspected aetiological agents. However, these agents, including CMV, HHV-6 and HPV are only found in a minority of the KS lesions (Huang et al. 1992; Kempf et al. 1995; Monini et al. 1996b). Following the discovery of KSHV (or HHV-8) in AIDS-KS biopsies (Chang et al. 1994; Moore & Chang, this issue), the vast majority of KS lesions from patients with AIDS and in the other epidemiological groups were shown to be positive for this novel virus (table 2) (Ambroziak et al. 1995; Boshoff et al. 1995b; Chang et al. 1996; Schalling et al. 1995). KSHV is now accepted to be the transmissible agent of KS.

KSHV DNA is found in all clinical stages of KS lesions (patch, plaque and nodular), and is generally absent in non-KS tissues from KS patients, in other vascular neoplasms and in other forms of skin tumours from immunosuppressed patients (IARC 1997). In peripheral blood mononuclear cells (PBMC), KSHV DNA can be detected by polymerase chain reaction (PCR) in approximately 50% of KS patients. Further, KSHV genome detection in PBMC of HIV-sero-positive individuals can predict who will subsequently develop KS (Moore et al. 1996; Whitby et al. 1995). Although one group reported the frequent detection of KSHV in the semen of healthy Italian donors (Monini et al. 1996a), KSHV is not detectable in semen donors in North America and the UK and only rarely in patients with KS (Howard et al. 1997; Pellett et al. 1999; Tasaka et al. 1996). Reports of KSHV detection in bone marrow stromal cells of multiple myeloma patients and in sarcoid tissues are also controversial (Di Alberti et al. 1997; Rettig et al. 1997) and most researchers now believe these are due to PCR contamination (Tarte et al. 1998; Whitby et al. 1997).

5. KAPOSI'S SARCOMA-ASSOCIATED HERPESVIRUS MOLECULAR STRAIN VARIATION

Open reading frame (ORF)-Kl is used to subtype KSHV (McGeoch & Davison 1999; Nicholas et al. 1998): subtypes A, B, C and D have been identified, and display between 15 and 30% amino-acid differences between their ORF-Kl-coding regions. These subtypes have close associations with the geographical and ethnic background of individuals (figure 1). Within these four subtypes, over 15 clades have now been described (Cook et al. 1999; Hayward 1999; Lacoste et al. 2000a; Meng et al. 1999; Poole et al. 1999; Zong et al. 1999). Subtype B is found almost exclusively in patients from Africa, subtype C in individuals from the Middle East and Mediterranean

Table 2. Detection of KSHV DNA by PCR in KS biopsies

(Data compiled from Ambroziak et al. 1995; Boshoff et al. 1995b; Buonaguro et al. 1996; Cathomas et al. 1996; Chang et al. 1994, 1996; Chuck et al. 1996; Dictor et al. 1996; Dupin et al. 1995; Gaidano et al. 1996; Jin et al. 1996; Lebbé et al. 1995, 1997; Luppi et al. 1996; McDonagh et al. 1996; Moore & Chang 1995; O'Neil et al. 1996; Schalling et al. 1995.)

type of lesion	no. positive/ no. tested	per cent positive
AIDS-KS	252/259	97
classic KS	160/175	91
iatrogenic KS	13/13	100
African endemic KS	71/80	89
HIV-negative gay men with KS	8/9	89
control tissues ^a	14/743	$1.8^{\rm b}$

^a Includes high-risks patients.

Europe, subtype A in western Europe and North America and subtype D has only so far been described in individuals from the Pacific Islands (Hayward 1999; Zong et al. 1999). A new subtype (E) has been described from South American indigenous people (Biggar et al. 2000). So far, no subtype appears to correlate with a specific disease entity or with a more aggressive course for KS. The unusually high genetic divergence identified in ORF-K1 reflects some unknown powerful biological selection process acting specifically on this immunoglobulin (Ig)receptor-like signal-transducing protein (Hayward 1999; McGeoch & Davison 1999). This could be related to evolving mechanisms of viral evasion from the immune system among different populations. Subtypes A and C are phylogenetically closest to each other, and subtype B more distant.

Subsequent to the discovery of KSHV, several new non-human primate rhadinoviruses have been isolated which have a closer phylogenetic relationship to KSHV than either Herpesvirus saimiri or EBV (see Damania & Desrosiers, this issue). These new members define two distinct rhadinovirus lineages. One lineage is closely related to KSHV and includes retroperitoneal fibromatosis herpesvirus (RFHV) (Rose et al. 1997), Chlorocebus rhadinovirus (ChRV) (Greensill et al. 2000) and chimpanzee (PanRHV1 and PanRHV2) and gorilla rhadinoherpesviruses (GorRHV1) (Lacoste et al. 2000b). The second lineage includes a rhesus monkey rhadinovirus (RRV) (Desrosiers et al. 1997) and ChRV2 (Greensill et al. 2000). The finding of two lineages of rhadinoviruses in nonhuman primates suggests that there might be an as yet unidentified RRV or ChRV2-type virus in humans. The possibility that KSHV was introduced from African apes to humans, rather than coevolved with Homo sapiens, is not excluded.

6. SERO-EPIDEMIOLOGY

(a) Serological assays

There are several KSHV serological assays currently available. The most widely used assays are based on the detection of latent or lytic antigens in KSHV-infected primary effusion lymphoma (PEL) cell lines, either by

 $^{^{\}rm b}p < 0.0001.$

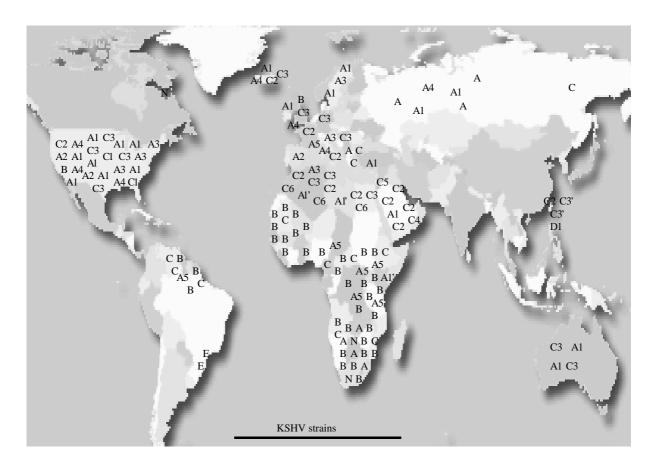


Figure 1. World distribution of KSHV strains and subtypes. The B and C subtypes described in North America are predominantly in those of African or Mediterranean origin. Data compiled from Cook et al. (1999), Davidovici et al. (2000), Hayward (1999), Lacoste et al. (2000a), Meng et al. (1999), Poole et al. (1999), Zong et al. (1999) and N. Wilder and C. Boshoff (unpublished data).

immunofluorescence (IF) (Gao et al. 1996a; Kedes et al. 1996; Simpson et al. 1996) or by enzyme-linked immunosorbent assay (Chatlynne et al. 1998). Assays have also been described which detect antibodies to recombinant KSHV latent and lytic proteins or synthetic peptides. Lytic proteins shown to be immunogenic include ORF 65 (Simpson et al. 1996), ORF 26 (Davis et al. 1997a), and ORF K8.1 (Chandran et al. 1998; Raab et al. 1998). The only latent antigen thus far to be used in recombinant assays is ORF 73, which is also the latent antigen detected in IF assays (Kedes et al. 1997b; Kellam et al. 1997; Rainbow et al. 1997). A study comparing various assays including recombinant proteins (ORFs 65, K8.1) and IF assays concluded that IF followed by confirmation with Western blot reactions with a panel of latent and lytic immunogenic antigens provide a reliable, sensitive and specific method to detect KSHV antibodies (Zhu et al. 1999).

(b) Sero-prevalence

(i) Northern Europe and North America

The sero-prevalence of KSHV in the different HIV-1 risk groups correlates with the incidence of KS: in the West, KSHV is found predominantly in HIV-positive gay men (Gao et al. 1996a; Kedes et al. 1996, 1997a; Lennette et al. 1996; Simpson et al. 1996) and not in HIV-positive heterosexuals, patients with haemophilia or intravenous drug users.

In a cohort of men in San Francisco, it was shown that KSHV infection is associated with the number of homosexual partners and correlates with a previous history of a sexually transmitted disease (e.g. gonorrhoea) and HIV infection (Martin et al. 1998), suggesting that KSHV is sexually transmitted. In a univariate analysis of gay men attending the sexually transmitted diseases clinic at St Thomas' Hospital in London, KSHV prevalence was associated with a history of sex with an American, suggesting that KSHV was perhaps first introduced into the gay communities in the epicentres of HIV in the USA, before spreading to Europe (Smith et al. 1999). In the Amsterdam gay men's cohort the risk of developing KS is higher when KSHV was acquired after HIV infection than vice versa (Renwick et al. 1998). This may be due to lower immune responses to primary KSHV infection in those who are HIV infected, with subsequently higher KSHV viral loads. There is also a suggestion of orogenital transmission of KSHV in this cohort (Dukers et al. 2000). In gay men in New York City and Washington DC the highest prevalence of KSHV infection occurred during the early 1980s and fell sharply thereafter (O'Brien et al. 1999).

(ii) Mediterranean Europe

The incidence of classic KS is significantly higher in Italy than in the UK or the US (Franceschi & Geddes 1995; Geddes *et al.* 1995), as is the prevalence of antibodies

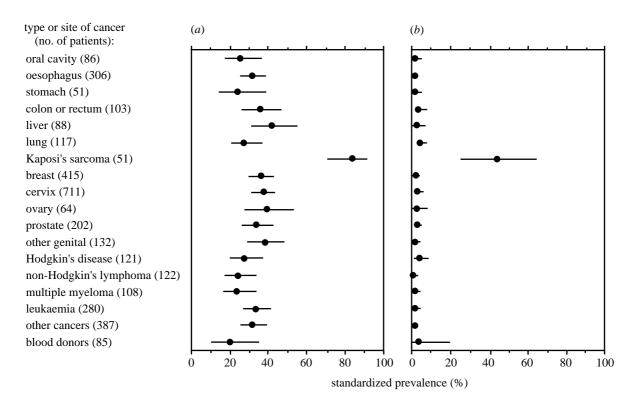


Figure 2. Antibodies to KSHV in 3400 black patients in South Africa with various types of cancer. (a) Sero-prevalence. (b) High-titre antibodies; presence of antibody titre is greater than 1:100 000. Adapted from Sitas et al. (1999).

to KSHV in blood donors (Cattani et al. 1999; Whitby et al. 1998). Furthermore, the incidence of classic KS in Italy shows considerable regional variation (Geddes et al. 1995) in parallel to KSHV prevalence (Calabro et al. 1998; Whitby et al. 1998). In addition, the geometric mean titre of anti-KSHV antibodies is highest in blood donors from the south, where the incidence of KS and the prevalence of KSHV is highest (Whitby et al. 1998). This finding is reminiscent of EBV infection, where high anti-EBV antibody titre correlates with an increased risk of developing Burkitt's lymphoma or nasopharyngeal carcinoma (de Thé et al. 1978a,b). The acquisition of KSHV in Italian children supports postnatal routes of transmission in families (Whitby et al. 2000).

(iii) Israel

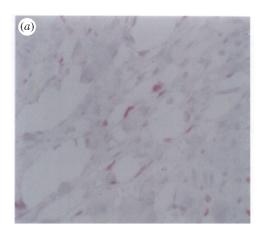
Various genetic diseases appear to cluster predominantly in either Ashkenazi Jews (eastern and northwestern Europe) or Sephardi Jews (North Africa and the Mediterranean). For example, Tay-Sachs syndrome is found almost exclusively among Ashkenazi Jews (Petersen et al. 1983), whereas familial Mediterranean fever is predominantly seen in non-Ashkenazi Jews, Arabs, Turks and Armenians (Ben-Chetrit & Levy 1998). The genetic mutations leading to these diseases would therefore appear to have been introduced after the two main arms of the diaspora separated. In contrast, the 185delAG mutation in BRCAl was originally described as a common mutation in Ashkenazi breast and ovarian cancer families (Tonin et al. 1996), but has since also been reported in Sephardi, Yemeni and Iranian Jews (Bar-Sade et al. 1998), suggesting a mutation prior to the dispersion of the Jewish people.

Classic KS is relatively common among both Sephardi and Ashkenazi Jews and the incidence of classic KS in Israel is among the highest in the developed world (Fenig et al. 1998; Zahger et al. 1993), compared with a low agestandardized incidence in the UK and USA (Biggar et al. 1984; Grulich et al. 1992) during a similar period (table 1). North African Jews (Sephardi) have a higher incidence of classic KS than Jews born in Israel or Europe (Iscovich et al. 1998; Marill et al. 1973). Thus KSHV or a genetic predisposition to KS appears to have been introduced into the Jewish population prior to the diaspora.

The sero-prevalence of KSHV among Israeli Jews is higher that that seen in the general populations of western Europe and North America (Davidovici et al. 2001) and the sero-prevalence of KSHV among the different Jewish groups correlates with the incidence of KS (Davidovici et al.). Place of birth and sero-status of spouse are the most important risk factors for adults to be infected. Furthermore, mother-to-child transmission is important in the acquisition of KSHV in Israel (Davidovici et al.). A relatively high prevalence of KSHV in Egyptian children (Andreoni et al. 1999) indicates similar routes of transmission in other Middle Eastern countries.

(iv) Africa

While endemic KS existed in parts of Africa long before the AIDS epidemic, AIDS–KS is now the most common tumour in many parts of Africa (table 1). In sub-Saharan Africa, where KS rates are relatively high among HIV-positive individuals, the prevalence of antibodies to KSHV is also higher than in North America and northern Europe (Ariyoshi et al. 1998; Gao et al. 1996; Olsen et al. 1998; Simpson et al. 1996; Sitas et al.



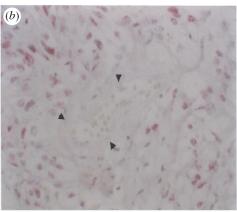


Figure 3. KSHV LANA-1 expression in early (a) and nodular (b) KS lesions. LANA-1 is only present in a minority of cells in early lesions, but in most spindle cells in advanced lesions. Normal mature endothelial cells are negative for KSHV

1999) and KSHV infection occurred in Africa before the epidemic of AIDS-KS (de Thé et al. 1999). In a study of KSHV serology in over 3400 black cancer patients (Sitas et al. 1999), only KS was correlated with prevalence and titre of KSHV antibodies (figure 2).

Acquisition of KSHV at an early age in Africa is likely because KS is seen in African children (Ziegler & Katongole-Mbidde 1996). Indeed, the prevalence of antibodies to KSHV increases steadily with age in Africa (Olsen et al. 1998; Rezza et al. 2000; Sitas et al. 1999) and this occurs before puberty (Bourboulia et al. 1998; Gessain et al. 1999; He et al. 1998; Mayama et al. 1998). This indicates that KSHV is not predominantly transmitted during sex, as with gay men in the West. Mother-to-child transmission and sibling-to-sibling transmission has been shown to occur in South Africa and in an Afro-Caribbean population living in French Guyana (Bourboulia et al. 1998; Plancoulaine et al. 2000). About one-third of KSHV-positive black mothers in South Africa transmit the virus to their children (Bourboulia et al. 1998). In South Africa there is significantly lower prevalence of anti-KSHV antibodies among whites than among blacks; in black cancer patients the sero-prevalence of KSHV declines with increasing education, suggesting that factors associated with poverty may contribute to the transmission of the virus (Sitas et al. 1999).

The lymph nodal form of KS seen in African children might be a manifestation of primary infection by KSHV, analogous to EBV-associated infectious mononucleosis. In support of this notion is the description of microscopical KS-type changes in the lymph node of an HIV-infected 43-year-old French patient at the time of presumed KSHV sero-conversion (Oksenhendler et al. 1998).

7. TRANSMISSION

Although one group reported the frequent detection of KSHV in the semen of healthy Italian donors (Monini et al. 1996a), in North America and the UK the current consensus is that KSHV is present only intermittently in the semen of patients with KS and sometimes in HIV-positive patients without KS, but only rarely in semen donors (Ambroziak et al. 1995; Corbellino et al. 1996a; Howard et al. 1997; Lin et al. 1998; Tasaka et al. 1996; Viviano et al. 1997). KSHV has also been reported in prostate biopsies of HIV-positive men with and without KS (Diamond et al. 1998; Staskus et al. 1997), so KSHV shedding into semen from prostate fluid is therefore a possible mode of transmission. Infectious virus is also found in the saliva of HIVpositive individuals (Koelle et al. 1997). In patients with classic KS, KSHV DNA was found in tonsillar swabs and in saliva (Cattani et al. 1999). Although studies in gay men indicate that KSHV is transmitted during sex and the risk of having KSHV increases with the number of sexual partners, the exact mode and route of transmission is not known. Semen and saliva are each possible routes of viral transmission, but their respective contribution to infection is unknown. The role of breast milk, saliva and other transmission routes for mother-to-child and sibling-tosibling transmission is also still unknown.

8. KAPOSI'S SARCOMA AND KAPOSI'S SARCOMA-**ASSOCIATED HERPESVIRUS INFECTION**

Four observations link KSHV causally to the aetiopathogenesis of KS, although none of these findings on their own is sufficient to support a causative role:

- (i) KSHV DNA is present, by PCR, in all epidemiological forms of KS, in all fresh biopsies tested and in the vast majority of paraffin-embedded material. However, KSHV DNA is rarely, if at all, detectable in other mesenchymal tumours (IARC 1997) (table 2).
- (ii) The detection of KSHV DNA by PCR in the peripheral blood of HIV-positive individuals predicts who might subsequently develop KS (Moore et al. 1996; Whitby et al. 1995), indicating that those at risk of KS have a higher viral load than those not at risk.
- (iii) Serological studies indicate that in general populations at risk of developing KS there is a higher prevalence of KSHV infection, so that the incidence of classic KS and AIDS-KS in different populations correlates broadly with the prevalence of virus infection in these populations.
- (iv) In advanced KS lesions, KSHV is latently expressed in nearly all the tumour (spindle) cells (Dupin et al. 1999; Sturzl et al. 1999). This is reminiscent of other viral cancers, e.g. EBV latent infection in posttransplant lymphoproliferative disease or HPV infection in cervical cancer.

To strengthen the molecular epidemiological association between KSHV and KS further, it was demonstrated by PCR in situ hybridization, RNA in situ hybridization and immunohistochemistry that KSHV is present in spindle cells in nearly all KS lesions (Boshoff et al. 1995a; Davis et al. 1997b; Dupin et al. 1999; Kellam et al. 1999; Li et al. 1996; Rainbow et al. 1997; Staskus et al. 1997; Sturzl et al. 1997) (figure 3). In early KS lesions, only a small proportion (<10%) of spindle cells are positive for KSHV, whereas VEGFR-3 is expressed by most cells (Dupin et al. 1999), indicating that paracrine mechanisms may be important in the initiation and progression of KS. In nodular lesions, more than 90% of the spindle cells contain KSHV latent infection, suggesting that KSHV latent proteins confer a growth advantage on infected cells (Dupin et al. 1999). KSHV is not present in other vascular tumours, including angiomas and angiosarcomas, and it is only rarely detectable in other forms of skin tumours (including squamous carcinomas and melanomas) in immunosuppressed patients (Adams et al. 1995; Boshoff et al. 1996; Lin et al. 1996; Uthman et al. 1996).

In culture, KSHV has been shown to infect macroand microvascular endothelium and to provide a growth advantage to these cells (Flore et al. 1998). With other oncoproteins like HPV E6 and E7, KSHV can transform human microvascular endothelium (Moses et al. 1999). Our current model is that KSHV, like EBV, persists in B cells as an episome expressing only a small number of latent proteins. Reactivation of the virus due to immunosuppression or local cytokine production leads to infection of endothelial precursors. In these precursors, KSHV latent proteins induce proliferation and block differentiation and apoptosis by anti-viral cellular immune responses. It is possible that local cytokine production stimulates neo-angiogenesis or neo-lymphangiogenesis and KSHV then infects these immature endothelial cells.

There is currently no evidence that KSHV infects endothelial cells in healthy individuals. Although the nature of the spindle cells in KS has been studied extensively there remains uncertainty regarding their exact origin. These cells usually express proteins that are found exclusively on endothelial cells. More recently, VEGFR-3/ Flt-4 was shown to be expressed by all the KSHV-positive spindle cells in KS lesions. VEGRF-3 is the receptor for the angiopromoting cytokines VEGF-C and VEGF-D. As VEGF-C promotes lymphangiogenesis and VEGFR-3 is highly expressed by mature lymphatic endothelia we propose that the KS tumour cell belongs to the lymphatic endothelium lineage of cells (precursors), rather than being de-differentiated mature microvascular endothelium (Dupin et al. 1999). The expression by all KS spindle cells of another marker related to lymphatic endothelium, podoplanin, supports this hypothesis (Weninger et al. 1999). In vitro, VEGF-C promotes a mitogenic and motogenic response of KS-derived spindle cells by activating the tyrosine phosphorylation of VEGFR-3 (Marchio et al. 1999).

The major viral latency-associated nuclear antigen (LANA-1 or LNA-1) encoded by ORF 73 appears to be expressed in virtually all KSHV-infected cells (Dupin et al. 1999; Parravicini et al. 2000; Katano et al. 2000). Using an antibody against LANA-1, Dupin et al. (1999) detected KSHV in nearly all spindle cells of well-developed KS lesions, but in only a few endothelial or spindle cells of early KS lesions, suggesting that paracrine mechanisms might be involved in the initiation of these lesions.

The histological, clinical and molecular features may be reconciled by invoking a model for KS in which early lesions are non-clonal proliferations of virus-infected endothelial or endothelial precursors and advanced disease represents oligoclonal or monoclonal neoplasms. This model is comparable with EBV-driven polyclonal lymphoproliferative disorders in immunodeficient individuals, which can progress to clonal lymphomas. The contribution to pathogenesis of lytically replicating virus in a subset of KS spindle cells has not been determined, although various lytic KSHV proteins trigger, at least in experimental systems, pathways involved in cell proliferation, anti-apoptosis, cell migration and angiogenesis (see Moore & Chang, this issue). In early lesions such a paracrine model could be involved in the pathogenesis of KS.

9. KAPOSI'S SARCOMA-ASSOCIATED HERPESVIRUS AND LYMPHOPROLIFERATION

(a) Body cavity-based primary effusion lymphoma

Two groups first recognized the unique aspects of some effusion-based lymphomas in patients with AIDS (Knowles et al. 1989; Walts et al. 1990). The cells in these cases were negative for most lineage-associated antigens, although a B-cell origin was indicated with clonal rearrangement of the Ig genes. Karcher et al. (1992) further demonstrated the distinctiveness of the syndrome, reporting a high prevalence of EBV and absence of *c-myc* rearrangements. Later, KSHV was specifically associated with primary effusion lymphoma (PEL), but not with any of the common types of Hodgkin's and non-Hodgkin's lymphomas (Cesarman et al. 1995; Gessain et al. 1997; Sitas et al. 1999).

PEL typically presents as malignant effusions in the visceral cavities usually without significant tumour mass or lymphadenopathy. These lymphomas occur predominantly in HIV-infected individuals with advanced stages of immunosuppression (Komanduri et al. 1996), but are occasionally seen in HIV-sero-negative patients (Nador et al. 1996; Strauchen 1997; Said 1996). As with KS, PELs are seen primarily in gay men (Jaffe 1996; Nador et al. 1996). PELs usually do not express surface B-cell antigens with the exception of CDl38/syndecan-1, a molecule selectively associated with late stages of B-cell differentiation (Gaidano et al. 1999). This finding, and frequent mutations in the 5' non-coding region of BCL-6, define PEL cells as preterminally differentiated, post-germinalcentre stage B cells (Gaidano et al. 1999) (table 3). It appears that KSHV-positive PEL cells lack many adhesion molecules and 'homing markers' present on other diffuse lymphomas: this may contribute to the peculiar effusion phenotype of these lymphomas and the usual lack of macroscopic lymph nodal involvement (Boshoff et al. 1998).

PEL cells consistently lack molecular defects commonly associated with neoplasia of mature B cells including activation of the proto-oncogenes c-myc, bcl-2, bcl-6, n-ras, and k-ras, as well as mutations of p53 (Nador et al. 1996). Southern blot analysis of PEL cells shows the presence of the KSHV genome in very high

Table 3. Comparison of KSHV-infected PEL cells and plasmablasts in multicentric Castleman's disease

(The term plasmablast is used to connote a medium-sized cell with a moderate amount of amphophilic cytoplasm and a large vesicular nucleus containing up to three prominent nucleoli. In contrast to an immunoblast, the cytoplasm contains abundant Ig (Dupin 2000).)

feature	PEL	plasmablastic multicentric Castleman's disease		
site morphology KSHV EBV cytoplasmic Ig expression Ig light chain CD30 CD138	body cavity, extranodal immunoblastic positive positive in majority absent monotypic κ or λ mRNA positive positive	lymph nodes, spleen plasmablastic positive negative high level of IgM monotypic λ Ig weakly positive not determined		
B-cell antigens mutation in Ig genes cellular origin	absent hypermutated in majority post-germinal-centre B cells	weak or absent absent naive $\operatorname{Ig} M\lambda$ -expressing B cells		

copy number (50-150 cell-1) compared with that seen in KS. Cell lines from PEL have been established (Arvanitakis et al. 1996; Cesarman et al. 1995; Renne et al. 1996). Some cell lines are only positive for KSHV; others are coinfected with EBV. In addition to cell lines established from lymphomatous effusions, the BCP-1 cell line was established from the peripheral blood of a patient with PEL suggesting that the malignant cells are present not only in the malignant effusions but may be also found disseminated in the peripheral blood (Boshoff et al. 1998). PEL cell lines express KSHV latent genes and can readily be induced by phorbol esters or butyrate into lytic replication (see Moore & Chang, this issue). Thus PEL cells have been of great use in defining KSHV gene expression and for the production of antigens, as well as for IF assay substrates (Gao et al. 1996b; Jenner et al. 2001; Kedes et al. 1996; Lennette et al. 1996).

A PEL-derived cell line, BC-3, was used to study the global expression of all known KSHV genes using gene expression microarray technology (figure 4). Cluster analysis, which arranges genes according to their expression profile, revealed a correlation between expression and assigned gene function that is consistent with the known stages of herpesvirus life cycles (Jenner et al. 2001). This study showed that non-induced PEL cells express a highly restricted number of KSHV genes (figure 4b) and these genes are presumably involved in maintaining the proliferation of these cells (see Moore & Chang, this issue).

KSHV has not yet been shown to transform or immortalize any B-cell type in vitro. However, the target cells used (i.e. mature B lymphocytes) may not be those that are susceptible to KSHV infection.

(b) Castleman's disease

Multicentric Castleman's disease (MCD) is a lymphoproliferative disorder of unknown aetiology and is associated with the development of secondary B-cell lymphoma (Castleman et al. 1956; Frizzera 1988). We now know that a subgroup of MCD is linked to KSHV. KSHV infection is found in nearly 100% of HIVassociated cases of MCD and 40-50% of HIV-negative cases (Chadburn et al. 1997; Corbellino et al. 1996b; Gessain et al. 1996; Soulier et al. 1995). HIV-sero-negative patients with KSHV-related MCD appear to experience a worse clinical course than KSHV-negative MCD and their disease is frequently complicated by autoimmune haemolytic anaemia and polyclonal gammopathies (Chadburn et al. 1997; Parravicini et al. 1997).

In MCD, KSHV is present in plasmablasts belonging to the B-cell lineage, which are not present in KSHVnegative MCD (Dupin et al. 1999; Katano et al. 2000). These plasmablasts localize mainly in the mantle zone of B-cell follicles. KSHV-positive MCD is therefore a distinct disease entity and designated as a plasmablastic variant of MCD (Dupin et al. 2000). Confluent clusters of KSHV-positive plasmablasts are also present in biopsies of plasmablastic MCD, indicating that isolated KSHV plasmablasts can progress to form foci of microlymphoma. KSHV is also present in all the tumour cells of plasmablastic lymphomas that develop in patients with the KSHV-positive plasmablastic variant of MCD (Dupin et al. 2000). The development of plasmablastic lymphoma therefore appears to represent a further evolution of this disorder (figure 5). Unlike KSHVpositive PEL cells, the plasmablasts in MCD are only positive for KSHV, and not for EBV (table 3).

The KSHV-positive plasmablasts express high levels of cytoplasmic IgM and show exclusively λ light chain restriction. In contrast, KSHV-negative, mature plasma cells in the interfollicular region, a prominent feature of MCD, are IgM negative and usually polytypic. KSHVpositive plasmablasts in MCD typically occur as isolated cells but may coalesce to form microscopic aggregates that show λ light chain restriction and therefore have been referred to as microlymphomas (Dupin et al. 2000). In some cases, KSHV-positive plasmablasts may form frank plasmablastic lymphoma. In all lesions in 25 patients studied so far, KSHV-positive plasmablasts uniformly express IgM\(\lambda\) (Dupin et al. 2000).

The finding that KSHV-positive plasmablasts in MCD are monotypic raises the question whether this is indicative of monoclonality or of preferential targeting of λ-expressing B cells by the virus (Dupin et al. 2000). It was recently shown that these monotypic cells are polyclonal as determined by Ig gene rearrangement (Du et al. 2001). KSHV infection therefore invokes a monotypic but



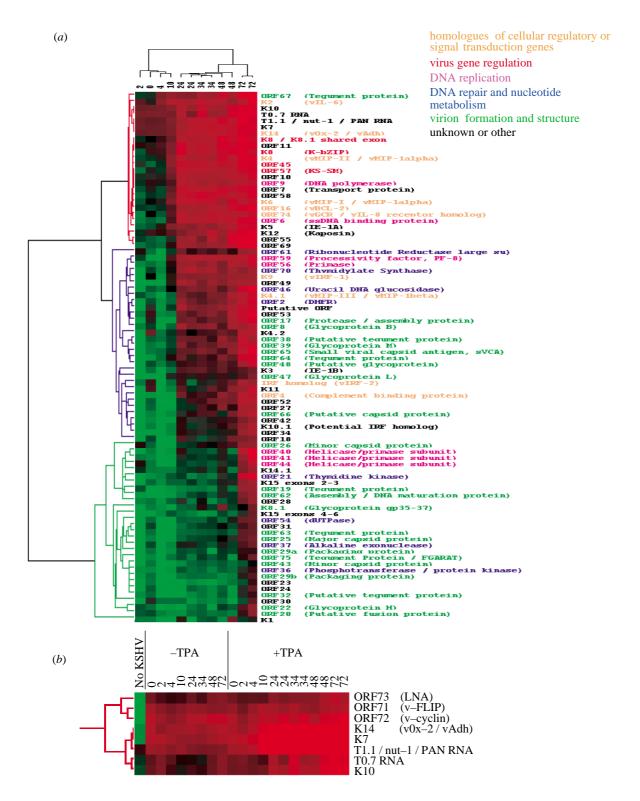


Figure 4. KSHV gene expression as determined by array technology and hierarchical clustering (from Jenner *et al.* 2001). (a) Hierarchical clustering of the genes and samples after the induction of lytic replication with TPA. The ORF and corresponding gene names are listed on the right-hand side and are colour-coded according to their putative function. (b) Expanded view of the cluster of genes whose expression is detectable in non-induced PEL cells.

polyclonal B-cell proliferation in MCD and in some cases KSHV-positive plasmablasts may form polyclonal or monoclonal microlymphomas or develop into frank monoclonal lymphoma (figure 5). These events (polyclonal to monoclonal evolution) are similar to those in

lymphoproliferative disorders caused by EBV in immunosuppressed hosts. However, EBV-associated lymphoproliferative disease, unlike KSHV-related lymphoproliferative disorders in MCD patients, tends to be of germinalcentre or post-germinal-centre origin.

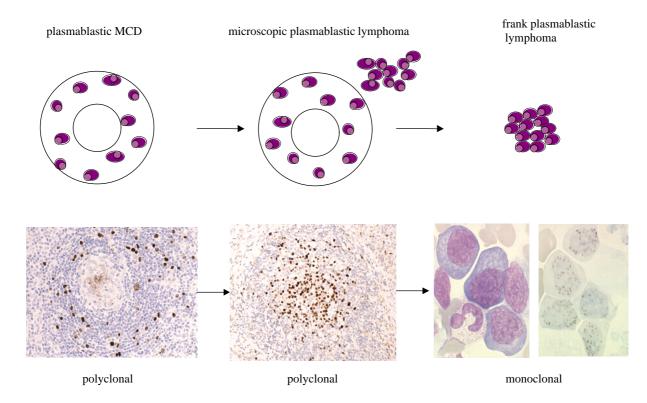


Figure 5. Model of KSHV in plasmablastic MCD. LANA-1 staining is shown in the bottom panel.

The reasons why KSHV is overwhelmingly associated with polyclonal λ light-chain-expressing B cells in MCD are not clear. In PEL, the tumour cells express functionally rearranged κ or λ light chain transcripts. In MCD, occasional KSHV-positive κ-expressing B cells are also seen. KSHV may naturally target both κ and λ light-chain-expressing B cells without bias, but λ cells may expand preferentially due to their intrinsic proliferative response to KSHV infection. The finding of κ light-chain-positive PEL does not argue against this hypothesis as the majority of PEL are also co-infected with EBV, which might override the differential response of κ and λ light-chain-expressing B cells to KSHV infection.

KSHV-positive plasmablasts do not harbour somatic mutations in the rearranged Ig gene, indicating that they originate from naive B cells despite their mature phenotype (table 3). These findings are consistent with their preferential localization in the mantle zones of B-cell follicles in MCD, and the lack of detectable follicular dendritic cell meshwork within the microlymphoma sites even though they are often adjacent to or partially replace B-cell follicles. Thus, KSHV may infect IgM⁺ naive B cells and drive these cells to differentiate into plasmablasts without undergoing the usual germinalcentre reaction, during which normal naive B cells mutate their rearranged Ig genes and differentiate into plasma or memory B cells. Like the plasmablasts in MCD, the plasmablastic lymphomas associated with MCD harbour non-mutated Ig genes and are derived from naive B cells. Two or even three independent KSHV-associated tumours (i.e. KS, PEL and MCD) in the same individual have been reported (Codish et al. 2000; Jones et al. 1998).

Viral IL-6 (vIL-6) is highly expressed in 10-15% of KSHV-positive plasmablasts (Du et al. 2001; Parravicini et al. 1998), and the IL-6 receptor is strongly expressed in the majority of KSHV-positive cells. Activation of the IL-6 signalling pathway may play an important role in driving KSHV-infected naive B cells to differentiate into plasmablasts and to develop various lymphoproliferative lesions. vIL-6 may directly stimulate both vIL-6-positive and -negative cells infected with KSHV by autocrine and paracrine mechanisms (see Moore & Chang, this issue). Elevated serum human IL-6 (hIL-6) has been demonstrated in patients with MCD and the level of hIL-6 correlated with the clinical presentation of the disease and high KSHV viral load in PBMC. Overproduction of hIL-6 is thought to be responsible for the systemic manifestation of MCD as blockage of IL-6 signalling by antibody can dramatically alleviate both clinical and histological presentations of the disease. Although it remains to be tested whether the elevated hIL-6 in MCD is the result of KSHV infection, it is noteworthy that hIL-6 is produced by PEL and promotes the growth of PEL cells in vitro and in vivo.

A role for HIV-1 in the pathogenesis of KS is supported by the observations that KS can rapidly resolve with highly active anti-retroviral therapy (HAART). Whether these responses are purely due to a restoration of cellular immunity against KSHV or because of the lower circulating HIV-1 load, is not yet known. In contrast, KSHVpositive MCD does not often resolve and can, despite HAART, progress to fatal lymphoma (Dupin et al. 2000; Zietz et al. 1999).

ORF 73 encodes LANA-1 of KSHV (Kedes et al. 1997b; Kellam et al. 1997; Rainbow et al. 1997). LANA-1 is expressed in all tumour cells in each of the KSHV-related

Table 4. Convergent evolution in the functions of certain DNA tumour viral nuclear phosphoproteins (Minus and plus signs denote does not bind to and binds to, respectively.)

	SV40 largeT antigen	papillomavirus		- KSHV	adenovirus	
		E6	E7	LANA-1	ElA	E1B
binds to pRB	+	_	_	+	+	_
binds to p53	+	+	_	+	_	+
binds to histone H1	5	_	+	+	?	?
maintains episome	5	+	+	+	+	_
transform primary rodent cells with a cellular oncogene	+	+	+	+	+	+

malignancies (figures 3 and 5). LANA-1 has been shown to maintain the KSHV episome, and tethers the viral genome to chromatin during mitosis via histone H1 (Ballestas *et al.* 1999; Cotter & Robertson 1999). LANA-1 interacts with the tumour-suppressor protein p53 and represses its transcriptional activity (Friborg *et al.* 1999). LANA-1 also binds to RING3 (Platt *et al.* 1999) and pRB (Radkov *et al.* 2000), two proteins involved in regulating E2F-dependent transcription. It is notable that other DNA tumour viral proteins that tether viral DNA to chromatin during mitosis also interfere with the p53 and pRb pathways (table 4).

10. CONCLUSIONS

Epidemiological evidence is overwhelming in support of KSHV as the central factor in the development of KS, the most common AIDS-related malignancy. The complex histology and expression pattern of KSHV proteins suggests that the role of the virus in KS pathogenesis is not straightforward and the model of KS tumorigenesis might not be like any other virally induced malignancy (Gallo 1998; Ganem 1996). The cell biology of KSHV also supports its role in the pathogenesis of a sub-type of MCD and of PEL.

The array of genes encoded by KSHV, some of which are unique to KSHV and others of which are shared only among the rhadinoviruses, has provided clues and directions in dissecting out the mechanisms of viral pathogenesis and oncogenesis. Investigation of these genes has begun to demonstrate their functional activity in cellular signalling and regulatory pathways (see Moore & Chang, this issue). Candidate oncogenes have been identified which may cause dysregulated proliferation or interfere with established tumour-suppressor pathways.

Like EBV, KSHV probably establishes a persistent infection which is normally controlled by the immune system and the number of KSHV-infected cells is under immunological control. When this immune control declines due to acquired or iatrogenic immunosuppression, the number of KSHV-infected cells increases with subsequent unchecked proliferation of virally infected cells and development of KSHV-related tumours. The introduction of HAART has led to a decline in the incidence of KS in AIDS patients and also to the resolution of KS in those already affected. This suggests that cellular immune responses, compromised in AIDS, but recovering after HAART, could be important in the control of

KSHV infection and in the development of KS. Remission of KS is also seen in organ-transplant patients upon cessation of immunosuppressive therapy. The rapid resolution of KS in some HIV-positive patients commencing HAART suggests that a small improvement in immunity might be important in disease control.

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